Claims:

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- A method of modifying a nucleic acid molecule comprising; contacting the nucleic acid molecule with a prokaryotic DNA repair ligase polypeptide.
- 2. A method according to claim 1 wherein the prokaryotic DNA repair ligase polypeptide comprises one or more of: a primase domain, a nuclease domain, and a ligase domain, said one or more domains sharing greater than 20% sequence identity with the corresponding domain sequence of Mt-Lig (CAB08492).
 - 3. A method according to claim 1 or claim 2 wherein the prokaryotic DNA repair ligase polypeptide shares greater than 20% sequence identity with the sequence of Mt-Lig (CAB08492).
- 20 4. A method according to any one of claims 1 to 3 wherein the prokaryotic DNA repair ligase polypeptide is Mt-Lig (CAB08492) or a variant or allele thereof.
- 5. A method according to any one of the preceding claims wherein the nucleic acid molecule and the Mt-Lig polypeptide are contacted in the presence of a prokaryotic Ku polypeptide.
 - 6. A method according to claim 5 wherein the prokaryotic Ku polypeptide shares greater than 20% sequence identity with the sequence of Mt-Ku (CAB08491).
 - 7. A method according to claim 6 wherein the prokaryotic Ku polypeptide is Mt-Ku (CAB08491) or an allele or variant thereof.
- 8. A method of ligating nucleic acid molecule ends comprising; contacting a first nucleic acid end and a second nucleic acid end with an prokaryotic DNA repair ligase polypeptide,

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wherein said first and said second nucleic acid ends are non-compatible.

- 9. A method according to claim 8 wherein said first and said second nucleic acid ends comprise non-complementary overhang regions.
- 10. A method according to claim 8 or claim 9 wherein the first end is on a first nucleic acid molecule and the second end is on asecond nucleic acid molecule.
 - 11. A method according to claim 10 wherein the first and second nucleic acid molecules are DNA.
- 15 12. A method according to claim 10 wherein the first nucleic acid molecule is DNA and the second nucleic acid molecule is RNA.
 - 13. A method according to claim 8 or claim 9 wherein the first and second ends are on the same nucleic acid molecule.
 - 14. A method according to any one of claims 8 to 13 comprising isolating and/or purifying the ligated nucleic acid molecule.
- 15. A method of labelling a nucleic acid molecule comprising;

 25 contacting a nucleic molecule having a first terminus with an prokaryotic DNA repair ligase polypeptide in the presence of labelled nucleotides.
- 16. A method according to claim 15 wherein the nucleotides are 30 NTPs.
 - 17. A method according to claim 15 wherein the nucleotides are dNTPs.
- 35 18. A method of filling in a single stranded gap in a double stranded nucleic acid molecule comprising;

contacting a double stranded nucleic acid molecule having a single stranded region with an prokaryotic DNA repair ligase polypeptide.

- 5 19. A method-according to claim -18 wherein said nucleic acid molecule and said prokaryotic DNA repair ligase polypeptide are contacted in the presence of NTPs.
- 20. A method according to claim 18 wherein said nucleic acid molecule and said prokaryotic DNA repair ligase polypeptide are contacted in the presence of dNTPs.
 - 21. A method of removing a single stranded overhang from the end of a nucleic acid molecule comprising;
- contacting said nucleic acid molecule with a prokaryotic DNA repair ligase polypeptide
 - 22. A method according to claim 21 wherein the prokaryotic DNA repair ligase polypeptide is an Mt-Lig polypeptide.

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- 23. A method according to claim 21 or claim 22 wherein said nucleic acid molecule is contacted in the presence of Mg^{2+} or Mn^{2+} .
- 24. A method of producing an RNA molecule comprising;
 25 contacting a prokaryotic DNA repair ligase polypeptide and a template DNA strand in the presence of NTPs.
 - 25. A method according to claim 24 wherein prokaryotic DNA repair ligase and template DNA are contacted in the presence of a primer oligonucleotide.
 - 26. A method of producing an DNA molecule comprising; contacting A prokaryotic DNA repair ligase polypeptide and a nucleic acid template in the presence of dNTPs and a primer oligonucleotide.
 - 27. A method according to claim 26 wherein the nucleic acid template is an RNA template.

- 29. A method according to any one of claims 8 to 28 wherein the prokaryotic DNA repair ligase polypeptide comprises one or more of: a primase domain, a nuclease domain, and a ligase domain, said one or more domains sharing greater than 20% sequence identity with the corresponding domain sequence of Mt-Lig (CAB08492).
- 30. A method according to any one of claims 8 to 29 wherein the prokaryotic DNA repair ligase polypeptide shares greater than 20% sequence identity with the sequence of Mt-Lig (CAB08492).

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- 31. A method according to any one of claims 8 to 30 wherein the prokaryotic DNA repair ligase polypeptide is Mt-Lig (CAB08492) or a variant or allele thereof.
- 32. A method according to any one of claims 8 to 31 wherein the nucleic acid molecule and the Mt-Lig polypeptide are contacted in the presence of a prokaryotic Ku polypeptide.
- 20 33. A method according to claim 32 wherein the prokaryotic Ku polypeptide shares greater than 20% sequence identity with the sequence of Mt-Ku (CAB08491).
- 34. A method according to claim 32 or claim 33 wherein the prokaryotic Ku polypeptide is Mt-Ku (CAB08491) or an allele or variant thereof.
 - 35. A kit comprising an isolated Mt-Lig polypeptide for use in a method according to any one of claims 1 to 34.
 - 36. A kit according to claim 35 comprising an isolated Mt-Ku polypeptide.
 - 37. A kit according to claim 35 or claim 36 comprising dNTPs.
 - 38. A kit according to claim 35 or claim 36 comprising NTPs.

- 39. A kit according to any one of claims 35 to 38 comprising one or more of buffers, stabilisers and excipients.
- --- 40.--A-method-of producing an prokaryotic_DNA_repair_polypeptide _______5 comprising;
 - (a) causing expression from nucleic acid which encodes a prokaryotic DNA repair polypeptide in a suitable expression system to produce the polypeptide recombinantly; and, testing the recombinantly produced polypeptide for prokaryotic DNA repair activity.
- 41. A method according to claim 40 wherein the recombinantly produced polypeptide is tested for one or more of: non-complementary end ligation activity, DNA dependent RNA primase activity, 3'-5' exonuclease activity, DNA and RNA dependent DNA polymerase activity, DNA dependent RNA polymerase activity, ATP dependent DNA and RNA ligase activity and DNA terminal transferase activity.
- 20 42. A method according to claim 39 or 40 wherein the prokaryotic DNA repair polypeptide is an Mt-Lig polypeptide or an allele or variant thereof.
- 43. A method according to any one of claims 39 to 41 comprising purifying said recombinantly produced polypeptide.